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(54) Title: A METHOD OF DETECTING AND DISCRIMINATING BETWEEN NUCLEIC ACID SEQUENCES (57) Abstract <p>The present invention is directed to a method for detecting the presence or absence of any specific target nucleic acid sequence contained in a sample. The target sequence can be present in the sample in a relatively pure form or as a component of a member of a mixture of different nucleic acids. The method of the invention utilizes a novel primer design. The sequence of the novel primer is composed of two portions, the 3' portion is a primer specific for the desired nucleic acid sequence and the 5' portion is complementary to preselected nucleic acid sequence. Extension of the 3' portion of the primer with a labeled deoxynucleosides triphosphate yields a labeled extension product if, but only if, the template includes the target sequence. The labeled extension product is detected by hybridization of the 5' portion to the preselected sequence. The preselected sequence is preferably bound to a solid support as one member of a grid having a group of sequences.</p>		

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REAGENTS COMPRISING CHIMERIC MOLECULES OF NUCLEIC ACIDS
AND NUCLEIC ACID ANALOGS

There are a series of compounds which have standard nucleic acid bases attached to nonstandard polymer backbones (these backbones can be nucleic acid like; eg. phosphorothioate- or methylphosphonate-linked deoxyribose; or they can be very different; eg. polyamide). If the interbase spacing is correct, these compounds are capable of base pairing reactions analogous to normal nucleic acids. Interest in these compounds currently centres on their use in either therapeutics or diagnostic imaging.

The chemical and hybridization properties of some of these compounds are particularly interesting. A number of novel and inventive uses of these compounds as tools in Molecular Biology (as opposed to therapeutic and diagnostic pharmaceuticals as above) are presented herein. The synthesis and use of non-standard backboned nucleic acids has been described in WO 92/20702 and WO 92/20703. More recently WO 93/12129 and WO 93/13121 described the synthesis and some uses of similar molecules.

In one aspect the present invention provides a chimeric nucleic acid/nucleic acid analogue molecule comprising a first portion, said first portion comprising a nonstandard backboned oligonucleotide having at least one amide linkage and a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond.

The preferred backbone for the nonstandard backboned oligonucleotide is polyamide. The oligonucleotide preferably comprises peptide nucleic acid (PNA).

Preferably, the nonstandard backboned oligonucleotide will contain at least two monomers, and more preferably three or more monomers. These may be

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monomers of PNA.

The chimeric molecules may be alternatively
backboned nucleic acids that have a few normally
backboned nucleotides at the 3' end. Such chimeric
5 molecules will combine the altered properties
associated with the nonstandard backbone with an
ability to prime DNA synthesis from a normal template-
bound 3' end. Some of these alternatively backboned
nucleic acids are capable of dsDNA strand invasion
10 below the target dsDNA melting temperature, somewhat
like ssDNA/RecA complexes (Science, 254, p1497,
(1991)).

Therefore new methods of sequencing are
possible. This is particularly the case for dsDNA. A
15 suitable chimeric molecule is mixed with the dsDNA to
be sequenced and then incubated with a DNA polymerase
and nucleoside triphosphates and appropriate
dideoxynucleoside triphosphate terminators. There
should be no need to denature the dsDNA to be
20 sequenced. This is a major advantage especially for
automated sequencing systems. In principle the 3' end
of the sequencing primer may have any chemical
functionality capable of acting as acceptor for the
formation of a phosphodiester bond. This may be an
25 -OH group on the non-standard backboned molecule
itself or a 3'-OH group if some normally backboned
nucleotide(s) are required. Only one normally
backboned nucleotide may be required but it is more
likely that a length of 2 to 12 bases, more preferably
30 3 to 7 bases may be required. Work described in
Biochemistry 29 1200-1207 (1990) shows that very short
conventionally backboned primers may be utilised.

In addition to chimeric molecules as
described herein, the invention also provides a method
35 of performing a primer extension reaction by the use

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of

- a) a target nucleic acid
 - b) a primer which is a chimeric nucleic acid/nucleic acid analogue molecule comprising a first
5 portion, said first portion comprising a nonstandard
backboned oligonucleotide and a second portion, said
second portion comprising an acceptor end which is a
chemical functionality capable of acting as acceptor
for the formation of a phosphodiester bond, said
10 chimeric molecule being capable of hybridising to part
of the target
 - c) a supply of nucleotides
- which method comprises mixing reagents a),
b) and c) in the presence of a chain extension enzyme
15 under conditions to allow the chimeric molecule to
hybridise to the target and extension of the chimeric
molecule at the acceptor end to occur, giving an
extension product.

- The invention also provides a method of
- 20 performing a chain termination reaction by the use of
 - a) a target nucleic acid
 - b) a chimeric nucleic acid/nucleic acid analogue
molecule comprising a first portion, said first portion
comprising a nonstandard backboned oligonucleotide and
25 a second portion, said second portion comprising an
acceptor end which is a chemical functionality capable
of acting as acceptor for the formation of a
phosphodiester bond, said chimeric molecule being
capable of hybridising to part of the target
 - 30 c) a supply of nucleotides
 - d) a chain termination agent
- which method comprises mixing reagents a), b)
c) and d) in the presence of a chain extension enzyme
under conditions to allow the chimeric molecule to
35 hybridise to the target and extension of the chimeric
molecule at the acceptor end to occur, so as to

produce terminated extension products, which terminated extension products are separated to allow part of the nucleotide sequence of the target nucleic acid to be determined.

5 A further aspect of the invention is a method of determining the nucleotide sequence of a target nucleic acid, which method comprises performing a chain termination method as described above, using a chain termination agent for each of the four different
10 nucleotides such that the nucleotide sequence of the target may be determined.

 The preferred chimeric molecules for use as primers in the methods described herein are chimeric molecules discussed above according to the invention.
15 Thus, the primers preferably have at least one amide linkage in the backbone of the nonstandard backboned oligonucleotide.

 Generally, a 6-mer is accepted as the smallest effective priming unit, although this can be
20 reduced to 3 or 4 if very low temperatures are used. The chimeric molecule primers described herein are preferably 6 or more base units in length.

 One or more of the reagents used in the various methods according to the invention may be
25 labelled in ways which are known in the art.

 There may be advantage in using an oligomer primer in which the majority (most or all) of the bases are pyrimidines. Such primers may prove to have enhanced strand invasion properties.

30 Whilst it will be appreciated that the use of chimeric molecules as herein described in Sanger chain-terminating sequencing is desirable, they may also be used in other related methods such as cycle sequencing.

35 The possibilities for using the chimeric molecules according to the invention and the

- 5 -

advantages these molecules provide in use also apply to other techniques including amplification techniques eg. PCR. It should therefore be possible to perform isothermal amplification of nucleic acids using the molecules herein described. Quantification of products produced by PCR reactions may also be improved by these molecules. In existing methods the dsDNA product of the PCR reaction is denatured and mixed with an immobilised capture probe. Some reannealing of the PCR product is inevitable and these molecules will not be captured by the immobilised probe leading to a loss of sensitivity. Use of PNA molecules, for example, in the capture step will eliminate the need to denature the PCR product leading to greater capture and increased sensitivity (PNA oligomers are peptide nucleic acid molecules comprising nucleic acid bases attached to a peptide backbone through a suitable linker and are described in detail in WO 92/20702 and WO 92/20703). The binding may also be stronger. The immobilisation of the capture probe may also be easier to achieve using conventional methods used for proteins and peptides.

Annealing of chimeric molecules at higher temperatures to provide increased specificity and greater sensitivity in primer extensions, PCR and sequencing reactions is expected.

A normally backboneed nucleotide or oligonucleotide could also be attached by its 3' end to the other end of the non standard backboneed nucleic acid so as to provide a normally backboneed 5' end and facilitate kinase labelling etc.

Non standard backboneed molecules could also be used in sequencing methods based on primer walking and subsequent developments by Studier (Science 11 December 1992 p 1787) . This original method makes use of sequence information near the terminus of a

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previously sequenced fragment of DNA to generate a new primer that will allow the next contiguous piece of DNA to be sequenced. This process can be repeated a number of times. One disadvantage of this approach, especially in genome sequencing projects, is the time required to determine and synthesise the next appropriate primer. This is slow and expensive.

The method developed by Studier uses a pre-synthesised pool of all possible different hexamers (approximately 4,000) of known sequence. These are combined to form 12mers or 18 mers to make the required primer. The hexamers can be ligated together on their complementary template or held together using a single stranded binding protein or just allowed to hybridise adjacent to each other.

If smaller primer units could be used then the number requiring synthesis would be much smaller. For example if 5mers could be used then approximately 1,000 would be required resulting in significant cost savings. Unfortunately, small oligonucleotides do not hybridise strongly. The strong hybridisation properties of, for example, PNA oligomers or other non-standard backboned molecules overcomes this problem. A combination of random PNA molecules and random standard oligonucleotides can be used to derive more efficient primer-walking strategies. The use of PNA-DNA chimeras which would hybridise more efficiently and also be primers for DNA polymerases is also possible.

The following list indicates some of the main uses for the chimeric molecules described herein. It will be clear to those skilled in the art that many other uses exist.

1. Primers for polymerases (using chimeric molecules with normally backboned nucleic acids giving the 3' end: priming could also occur from strand invasion

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- complexes formed by reaction with dsDNA targets, a strand displacing polymerase would be used for the extension). This can also be extended to random primer labelling of dsDNA molecules using chimeric primers and strand invasion.
- 5 2. In situ hybridization with no need to denature the sample.
 3. Primers/Probes that do not require dsDNA denaturation.
 - 10 4. Improved sequencing by hybridization with short oligos.
 5. Double strand sequencing without denaturation.
 6. Isothermal PCR and related PCR alternatives.
 7. Random primer labelling without denaturation.
 - 15 8. Better probes for primer extension mapping procedures.
 9. New labels and probes for multiplex sequencing.
 10. Novel adapters/linkers for cDNA cloning.

20

TERMINOLOGY:

By nucleic acid(s), we mean either DNA or RNA of any chain length which can be either wholly or partially single or double stranded unless otherwise specified.

25

By "normally backboned", we refer to phosphodiester linked deoxyribose (for DNA) or phosphodiester linked ribose (for RNA) as the backbone to which the base residues (A, C, G and T for DNA and A, C, G and U for RNA) are linked.

30

By "nonstandard backboned", we refer to any polymers other than the normal phosphodiester linked deoxyribose (for DNA) or phosphodiester linked ribose (for RNA) as the backbone to which the base residues (A, C, G and T for DNA and A, C, G and U for RNA) are linked. Unless otherwise stated, the only requirement

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for the nonstandard backbone polymer is that the interbase spacings are suitable for the formation of appropriate hydrogen bonds (Watson and Crick or triple helical or Hoogsteen type) with a normally backbone nucleic acid target. Examples of nonstandard backbones are phosphorothioate linked deoxyribose, phosphorothioate linked ribose, methylphosphonate linked deoxyribose, methylphosphonate linked ribose and polyamide. It will be immediately obvious to one skilled in the art that there are many other possible backbone polymers allowing the correct interbase spacing and that this allows for a number of different chemical and physical properties specific to the backbone moiety to be exploited whilst preserving the ability to bind to a complementary nucleic acid base sequence by hydrogen bonding.

It is also to be understood herein that chimeric molecules comprising nonstandard backbone nucleic acids with normally backbone nucleic acid ends, for ligation, priming, labelling and other such applications known to those skilled in the art, are also possible. Such chimeric molecules are to be included in the Claims wherever the term "nonstandard backbone" is used unless otherwise specified. In chimeric molecules the normally backbone sequence may be directly linked to the non-standard backbone sequence. It is also possible to include a small linker group between the two sequences.

By hybridization, we mean the sequence specific binding between a probe (with A, C, G and T residues or A, C, G and U residues attached to a normally backbone or nonstandard backbone polymer as specified) and a target nucleic acid. For a wholly or partially double stranded target, the sequence specific binding may also occur in a double stranded region by a process referred to herein as "strand

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invasion". Strand invasion is where the sequence specific binding of probe occurs under conditions in which the target strands do not normally separate from each other (for example at temperatures below the melting temperature of the target in a given solvent at a given ionic strength). Strand invasion does not normally occur with normally backboned probes. The application of this strand invading property of some nonstandard backboned nucleic acid probes to improve existing and create novel Molecular Biology applications is a major inventive step.

Figures 1 to 5 show HPLC traces for the DNA and PNA/DNA molecules synthesised in Example 6:

Figures 1a and 1b - 15-mers of DNA and PNA/DNA, respectively;

Figures 2a and 2b - 12-mers of DNA and PNA/DNA, respectively;

Figures 3a and 3b - 9-mers of DNA and PNA/DNA, respectively;

Figures 4a and 4b - 7-mers of DNA and PNA/DNA, respectively;

Figures 5a and 5b - 5-mers of DNA and PNA/DNA, respectively.

EXAMPLES

Example 1

Synthesis of chimeric molecule with single normal backboned base

0.5mg of PNA₁₅₄-NH₂ (H-CAT CTA GTG A-LysNH₂), synthesised according to the methods disclosed in WO 92/20702 and WO 92/20703, was mixed with 0.1mg 5'-amino

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thymidine in 0.02ml 33%DMSO, 100 mM Tris-HCl pH 7.4.
The reaction was started by the addition of 0.05mg
subaric acid bis(N-hydroxysuccinimide) ester and
incubated at room temperature for 24 hours.

5

Example 2Synthesis of chimeric molecule with longer standard
backboned sequence

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0.5mg $5'$ H₂NTCG CAC TGC ATC $3'$ (standard backbone) in
0.25ml 100mM Tris-HCl pH 7.4 was mixed with 0.5mg
subaric acid bis(N-hydroxysuccinimide) ester in DMSO
for 5 minutes. The reactants were purified by gel
15 filtration, freeze dried and resuspended in 0.1ml 100mM
Tris-HCl pH 7.4. 0.5mg PNA₁₅₄ in 0.025ml Tris-HCl pH
7.4 was added and incubated for 24 hours at room
temperature.

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Example 3PNA-DNA PRIMER EXTENSION ASSAY - GENERALISED PROTOCOLAnnealing conditions

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The PNA-DNA primer synthesised as in
examples 1 and 2 is diluted, 0 , 1/10 1/100 and 1/1000
in 50mM tris, pH 7.5 containing 50mM NaCl and 7mM
MgCl₂.

30

The control DNA 11mer primer is diluted to
10pmole/ μ l in the same buffer.

The PNA-DNA primer and the control primer are mixed
with equal volumes of 2pmole/ μ l template oligo to give
1pmole/ μ l template concentration.

35

Annealing is performed by boiling the
mixtures for 3 minutes and then leaving the solutions

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to cool to room temperature over a period of approximately 1 hour .

Primer Extension Assay

5

The extension reactions are carried out containing the PNA-DNA or the DNA primed template with 200µM dATP, dGTP, dTTP and 20µCi of alpha ³²P dCTP with exonuclease free Klenow polymerase.

10

The reactions are incubated at 37°C for 20 minutes then dCTP is added to a final concentration of 200µM. The reactions are incubated for a further 10 minutes then terminated by the addition of EDTA. Free nucleotide is removed by spin column centrifugation.

15

The results can be analysed by conventional denaturing polyacrylamide electrophoresis.

20

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30

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Example 4

The synthesis of the thymine PNA monomer is outlined in scheme 1.

All solvents were of HPLC grade; DCM and pyridine was distilled over CaH_2 ; MeOH was distilled over Mg and I_2 ; EtOH was distilled over CaO then Mg and I_2 ; DMF was of peptide synthesis grade and was not further purified; anhydrous acetonitrile was purchased from Applied Biosystems Inc. (ABI). All other chemicals were supplied by Aldrich.

^1H and ^{13}C n.m.r. were recorded on a Bruker 250AC and Bruker 200WP spectrometer. Positive ion Fast Atom Bombardment (FAB) mass spectra were recorded on a Kratos MS50TC spectrometer using a thioglycerol matrix. Oligonucleotide synthesis was carried out on an ABI 394 DNA synthesiser. PNA/DNA synthesis (detritylation and cleavage from the solid support) was carried out on an ABI 380B DNA synthesiser.

Flash chromatography was carried out using silica gel 60 (Fluka). Thin layer chromatography (tlc) was carried out on aluminium sheets, silica 60 F254, 0.2mm layer (Merck) using the following solvent systems; A=nBuOH/AcOH/ H_2O (3:1:1 v:v), B=DCM/EtOAc (1:1 v/v), C=DCM/MeOH (9/1 v/v). Products were visualised using ninhydrin (1% w/v in EtOH), with heating for 5 min. ,short wave UV lamp(254nm) or via iodine oxidation.

1-Thyminyllactic acid (I)

Thymine (10.0g, 79.3 mmols) was dissolved in water (50ml) containing KOH (17.1g, 0.30 moles) at 40°C . A solution of $\text{BrCH}_2\text{CO}_2\text{H}$ (16.5g, 1.5 eq.) in water (25ml) was added dropwise over 30 min. and the reaction mixture heated at 40°C for 2h. The solution was cooled to room temperature, the pH adjusted to 5.5 (c. HCl) then stored at -4°C for 2h. The precipitate formed was removed by filtration and the filtrate adjusted to pH 2 (c. HCl). The precipitate formed was isolated by filtration and dried over P_2O_5 (12.5g, 86%). $R_f=0.1(\text{A})$, FAB MS 185 ($\text{M}+1$) $^+$, ^1H n.m.r. (200MHz, DMSO); 1.74(s,3H,Me), 4.36(s,2H, CH_2), 7.48(s,1H,Ar-H), 11.4(s,1H,NH). ^{13}C n.m.r. (50MHz, DMSO); 12.1(Me), 48.7(CH_2), 108.6(C_5), 142.1(C_6), 151.2(C_2), 164.7(C_4), 169.9(CO_2H).

3-t-butoxycarbonylamino-1,2-propanediol (II)

3-amino-1,2-propanediol (10g, 0.11moles) was dissolved in water (250mL). The solution was cooled to 0°C and Boc anhydride (25g, 0.12 moles) added in one portion. The reaction mixture was brought to room temperature and the pH maintained at 10.5 using NaOH (2M) until the pH remained constant. The reaction mixture was stirred for a further 1 hour at room temperature then EtOAc (200mL) added and the pH adjusted to 2.5 using HCl (4M) at 0°C. The phases were separated and the aqueous phase extracted with EtOAc (10x150mL). The combined organic phases were dried (MgSO₄) then evaporated *in vacuo* to yield V as an oil. The product solidified upon freezing (20.4g, 97%). $R_f=0.75(A), 0.60(B)$, FAB MS 192 (M+1)⁺, ¹H n.m.r. (200MHz, CDCl₃); 1.39(s,9H,Boc Me), 3.14(broad, 2H, CH₂), 3.52(m,2H,CH₂), 3.68(m,1H,CH), 4.75(s,2H,2xOH), 5.52(m,1H,NH). ¹³C n.m.r. (50MHz, CDCl₃); 28.17(Me), 42.55(NCH₂), 63.48(CH₂), 71.08(CH), 79.76(quaternary C), 157.1(C=O).

t-butoxycarbonylaminoacetaldehyde (III)

The diol V (5g, 26.1mmoles) was dissolved in water (50mL) and NaIO₄ (6.8g, 1.2eq.) added. The reaction mixture was stirred at room temperature for 3h., filtered, then the aqueous solution extracted with DCM (5x100mL). The combined organic phases were dried (MgSO₄) then evaporated to dryness *in vacuo* to yield the aldehyde as a colourless oil (3.9g, 94%). The product was used without further purification in the next step. $R_f=0.7(B), 0.15(A)$; ¹H n.m.r. (200MHz, CDCl₃); 1.38(s,9H,Boc), 3.98(s,2H,CH₂), 5.33(b,1H,NH), 9.56(s,1H,CHO). ¹³C n.m.r. (50MHz, CDCl₃); 28.0 (Me), 51.1(CH₂), 79.9(qC), 155.6(Boc C=O), 197.4(CHO).

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N-(t-butoxycarbonylaminoethyl)-glycine ethyl ester (IV)

The aldehyde III (5.13g, 32.2 mmol) was dissolved in MeOH (100ml) and added to a mixture of ethyl glycinate (HCl salt, 11.2g, 2.5eq.) and NaBH₃CN (2.02g, 1eq.). The reaction mixture was stirred at ambient temperature for 16h. then the solvent removed *in vacuo*. The residue was dissolved in water (100ml) and the pH adjusted to 8.0 using NaOH (2M). The aqueous phase was extracted with DCM (5x150ml), the combined organic fractions dried (MgSO₄), then evaporated *in vacuo* to an oil which was purified by Kugelrohr distillation (6.92g, 87%) R_f=0.55(B), 0.15(C),
¹Hn.m.r. (250MHz,CDCl₃); 1.09(t,3H,CH₃), 1.25(s,9H,BocMe),2.53(t,2H,CH₂), 2.90(b,1H,NH), 3.46(t,2H,CH₂NH), 3.54(s,2H,CH₂),4.00(q,2H,CH₂CH₃),5.46(b,1H,carbamateNH).
¹³Cn.m.r.(50MHz,CDCl₃); 13.69(CH₃CH₂), 27.91(BocMe), 39.6, 42.6, 48.27, 60.36(all CH₂), 78.58(quat. C), 156.1(BocC=O), 172.4(C=O).

N-(t-butoxycarbonylaminoethyl)-N-(thyminyllacetyl)-glycine-ethyl ester (V)

Bocaminoethylglycine ethyl ester (IV, 2.75g, 11mmol) was dissolved in DMF (12mL) and thyminyllacetic acid (I, 2.05g, 1eq.) was added. On dissolution of the acid DCM (10mL) was added, the reaction mixture cooled to 0°C and DCC (2.49g, 1.2eq.) added. The reaction was stirred at 0°C for 1h. then for a further 2h. at room temperature. The precipitated DCU was removed by filtration, washed with DCM (2x30mL) and a further volume of DCM (150mL) added. The combined organic phases were washed with NaHCO₃ (0.5M, 3x100mL), citric acid (10%w/v, 2x100mL), and NaCl solution (sat., 1x100mL), dried (MgSO₄) then evaporated to dryness *in vacuo* to yield a brown oil. The crude product was purified by silica gel chromatography eluting with DCM/MeOH(1-3%) to yield the product as a white foam (1.86g, 41%), R_f=0.55(A),0.15(C) FAB MS 413 (M+1)⁺, ¹H n.m.r. (250MHz,CDCl₃); (some of the signals are split into major and minor peaks due to restricted rotation around the secondary amide bond.) 1.18(t,3H,MeCH₂), 1.35(s,9H,BocMe), 1.81(s,3H,TMe), 3.25(m,2H,CH₂), 3.50(m,2H,CH₂),

4.00(s,2H,NCH₂CO₂Et), 4.39(s,mi,CH₂CON), 4.53(s,mj,CH₂CON), 5.76(b,1H,BocHN), 6.99(s,mj,Ar-H), 7.0(s,mi,Ar-H), 10.0(s,mj,NH), 10.05(s,mi,NH). ¹³C n.m.r. (50MHz,CDCl₃); 12.1(CH₃CH₂), 13.8(T-Me), 28.14(BocMe), 38.4, 47.6, 48.4, 48.7, 61.4(all CH₂), 79.6(quat.C), 140.84(C6), 151.0, 155.9, 164.4, 167.2, 169.4(all C=O).

N-(Monomethoxytritylaminoethyl)-N-(thiminylacetyl)-glycine-ethyl ester (VI)

1-(Bocaminoethylglycine)-thymine ethyl ester (VII, 2.0g, 4.85mmoles) was dissolved in TFA (10mL). The reaction mixture was evaporated to dryness *in vacuo* then the residue coevaporated with toluene (5x10ml) then pyridine (2x10mL). Monomethoxytrityl chloride (1.65g, 1.1eq.), 4-pyrrolidinylpyridine (25mg) and pyridine (10mL) were added and the solvent was removed *in vacuo* once more. Pyridine (10mL) was added and the reaction mixture was stirred for 16h. Water (30mL) was added and the reaction mixture extracted with DCM (3x70mL). The combined organic phases were dried (MgSO₄), then evaporated to dryness *in vacuo* to yield an oil. The product was isolated by column chromatography using DCM/MeOH(0-3%) as the eluant (2.55g, 90%). The SiO₂ was pre-equilibrated with DCM containing NEt₃ (1%v/v) to prevent detritylation. R_f=0.2(A), 0.1(A); FAB MS 585 (M+1)⁺, ¹H n.m.r. (250MHz, CDCl₃); (some of the signals were split into major and minor peaks due to restricted rotation about the secondary amide bond) 1.18(t,3H,MeCH₂), 1.90(s,3H,T-Me), 2.38(m,2H,CH₂NCO), 3.52(m,2H,TrNH-CH₂), 3.76(s,3H,OMe), 3.90(s,mj,CH₂CON), 4.02(s,mi,CH₂CON), 4.10(q,2H,CH₂Me), 4.25(s,mj,CH₂CO₂Et), 4.31(s,mi,CH₂CO₂Et), 6.82(s,mi,H6), 6.85(s,mj,H6), 7.2-7.4(m,Ar-H). ¹³C n.m.r.(50MHz, CDCl₃); 12.2(CH₃CH₂), 13.9(T-Me), 41.7, 48.0, 48.5, 50.0, 61.3(all CH₂), 55.0(OMe), 110(CPh₃), 123-128(m,Ar-H), 137.4(quatC), 140.8(C6), 145.6(quatC), 149.5(quatC), 151.0(COMe), 157.8, 164.1, 167.3, 168.8(allC=O).

N-(Monomethoxytritylaminoethyl)-N-(thyminylacetyl)-glycine-(VII)

The ester XI (1.34g, 2.3mmoles) was dissolved in MeOH (60mL) and NaOH (2M, 40mL) added. The reaction mixture was stirred at room temperature for 2h. then DOWEX (pyridinium form) was added until a pH of 7 was obtained. The suspension was filtered and the resin washed with MeOH (3x50mL). The filtrate was evaporated to dryness *in vacuo* to yield XI (1.4g, 96%). $R_f=0.05(B), 0(C)$; 1H n.m.r. (200MHz, $CDCl_3$); (some of the signals were split into major and minor peaks due to restricted rotation about the secondary amide bond) 1.75(s, 3H, T-Me), 2.21(m, 2H, CH_2), 3.49(m, 2H, CH_2), 3.76(s, 3H, OMe), 3.85(s, mj, CH_2CON), 4.05(s, mi, CH_2CON), 4.52(s, mj, CH_2CO_2H), 4.85(s, mi, CH_2CO_2H), 6.85(s, mj, H6), 6.90(s, mi, H6), 7.2-7.5(m, Ar-H). ^{13}C n.m.r. (50MHz, $CDCl_3$); 12.08(T-Me), 33.48, 42.15, 48.23, 48.38 (all CH_2), 55.07(OMe), 113.2(CPh₃), 126-131(m, Ar-H), 137.4, 146.0, 149.8 (all quat C), 150.8(COMe), 158.9, 164.4, 168.3, 173.3(C=O).

Example 5

The nucleoside 5'-monomethoxytritylamino-5'-deoxy thymidine phosphoramidite was synthesised as shown in scheme 2 and was used as the initial linker in the PNA DNA chimeric molecule.

5'-azido-5'-deoxy thymidine (VIII)

Thymidine (14.0g, 58.0mmoles) was stirred in DMF (175ml) and PPh_3 (18.2g, 1.2eq.), NaN_3 (11.3g, 3eq.), and CBr_4 (23.0g, 1.2eq.) added at 0°C. The reaction mixture was stirred at 0°C for 1h. then ambient temperature for 16h. The reaction mixture was poured onto $NaHCO_3$ (0.5M, 350ml) and extracted with DCM (6x150ml). The combined organic phases were dried ($MgSO_4$) then evaporated to an oil which was purified by SiO_2 chromatography (0-4% MeOH/DCM, 7.73g, 50%). $R_f=0.1(C), 0.8(A)$, FABMS (M+1) 268, 1H n.m.r. (200MHz, D_6 -DMSO) 1.78(s, 3H, T- CH_3), 2.10(m, 1H, 2''H), 2.26(b, 1H, 2'H), 3.56(d, 2H, 5''H), 3.85(m, 1H, 4'H), 4.20(m, 1H, 3'H), 5.42(d, 1H, 3'OH), 6.21(t, 1H, 1'H)

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7.5(s,1H,6H), 11.34(s,1H,NH). ^{13}C n.m.r.(50MHz, D6-DMSO); 12.2(CH₃), 33.9(CH₂), 51.8(CH₂), 70.9,72.1(CH), 136.2(C6), 150.6, 163.8(C=O).

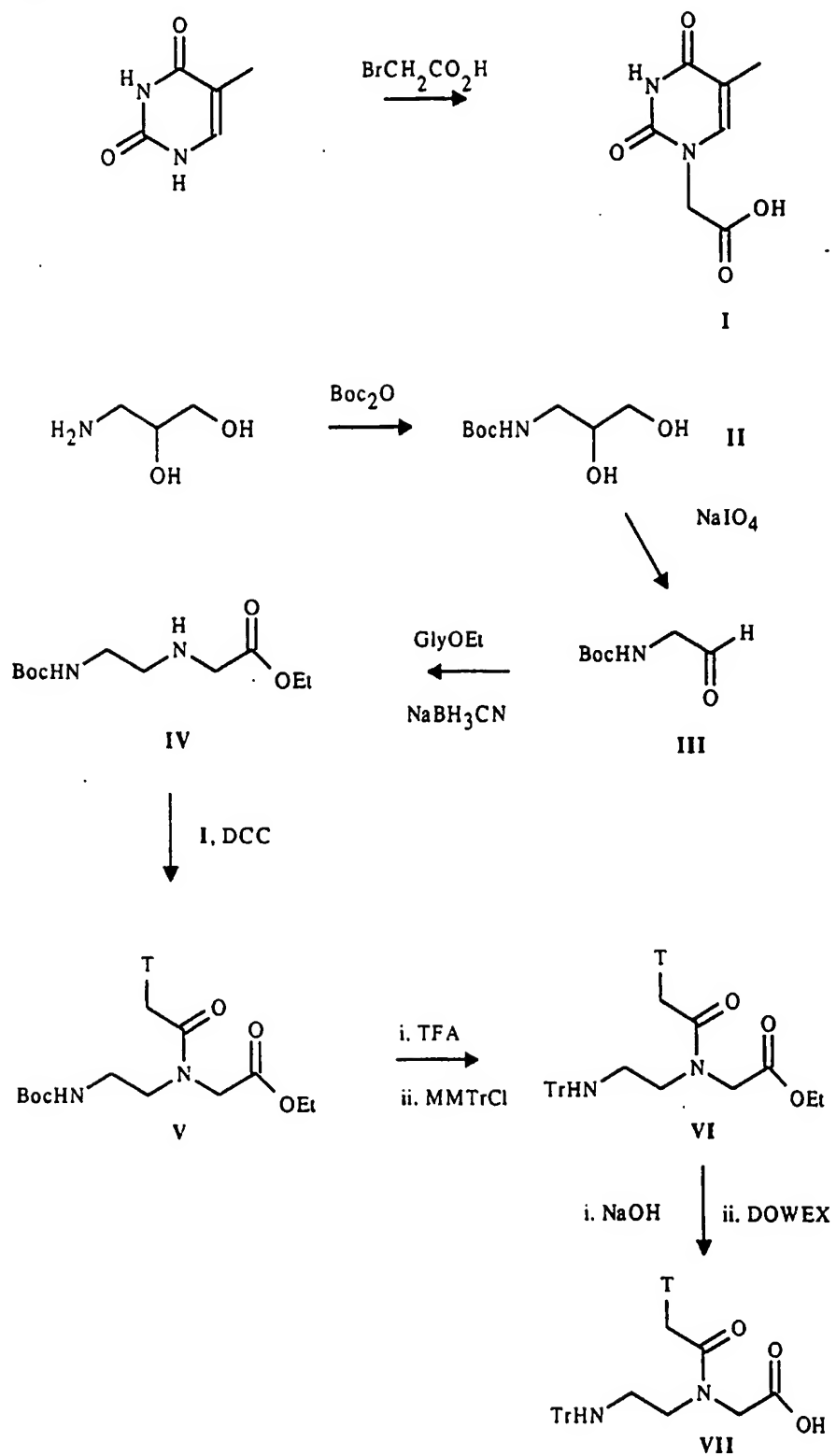
5'-N-(4-Methoxytrityl)amino-5'-deoxythymidine (X)

To the azido compound (7.73g, 29mmoles) in pyridine (100ml) was added PPh₃ (15.2g, 1.2 eq.). The reaction mixture was stirred at room temperature for 16h. then water (20ml) added. After stirring for 1h. the reaction mixture was poured onto water (300ml) and stirred for a further 16h. The suspension was filtered and the aqueous phase washed with EtOAc (3x150ml). The 5'-amino-5'-deoxy-thymidine was isolated by freeze drying (4.1g, 58%).

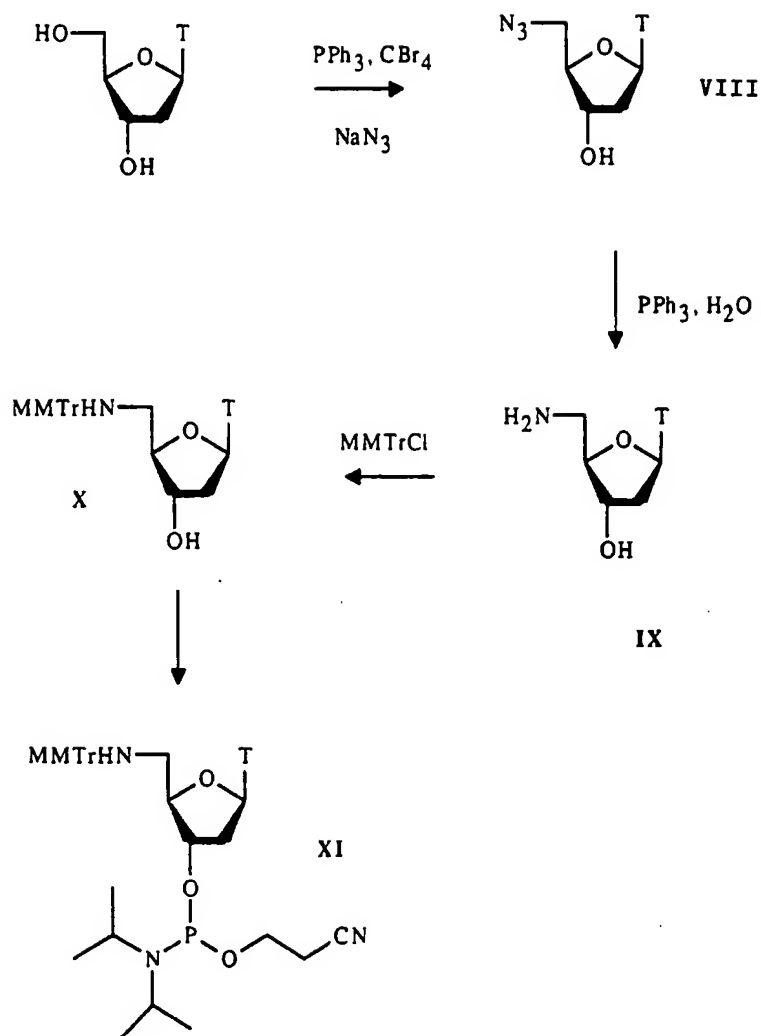
A fraction of the 5'-amino-5'-deoxy-thymidine (1.34g, 5.55 mmole) was coevaporated with pyridine (2x20ml) then dissolved in pyridine (40ml). 4-pyrrolidinopyridine (30mg) and p-anisoyldiphenylmethyl chloride (2.05g, 1.2eq.) were added, the reaction mixture stirred at room temperature for 16h. then poured onto NaHCO₃ solution (0.5M,30ml). The aqueous phase was extracted with DCM (3x50ml), the combined organic phases dried (MgSO₄) then evaporated to dryness *in vacuo*. The resultant oil was purified by column chromatography (SiO₂ preequilibrated with 1% NEt₃ in DCM eluting with 0-3% MeOH/DCM). (2.02g,71%) R_f=0.5(C), 0.2(B). FABMS (M+1)=514 ^1H n.m.r. (200MHz, D6-DMSO); 1.69(d,3H,CH₃), 2.01-2.34(m,4H,2'H,5'H), 2.64(t,1H,5'NH), 3.72(s,3H,OMe), 3.83(m,1H,4'H), 4.19(m,1H,3'H), 5.22(d,1H,3'-OH), 6.14(t,1H,1'H), 6.84 (d,2H,OMe), 6.87-7.43(m,13H,aromatic), 11.29(s,1H,NH). ^{13}C n.m.r.(50MHz, D6-DMSO); 12.2(CH₃), 24.9(CH₂), 40.0(CH₂), 54.8(OMe), 71.6(CH), 84.0(CH), 85.9(CH), 112.8-129.4(aromatic CH), 134.9, 137.4, 145.6, 147.8, 157.5(all qC), 150.4(C=O), 164.1(C=O).

**5'-N-(4-Methoxytrityl)amino-5'-deoxythymidine-3'-O-(2-cyanoethyl,
N,N-diisopropylamino)phosphite (XI)**

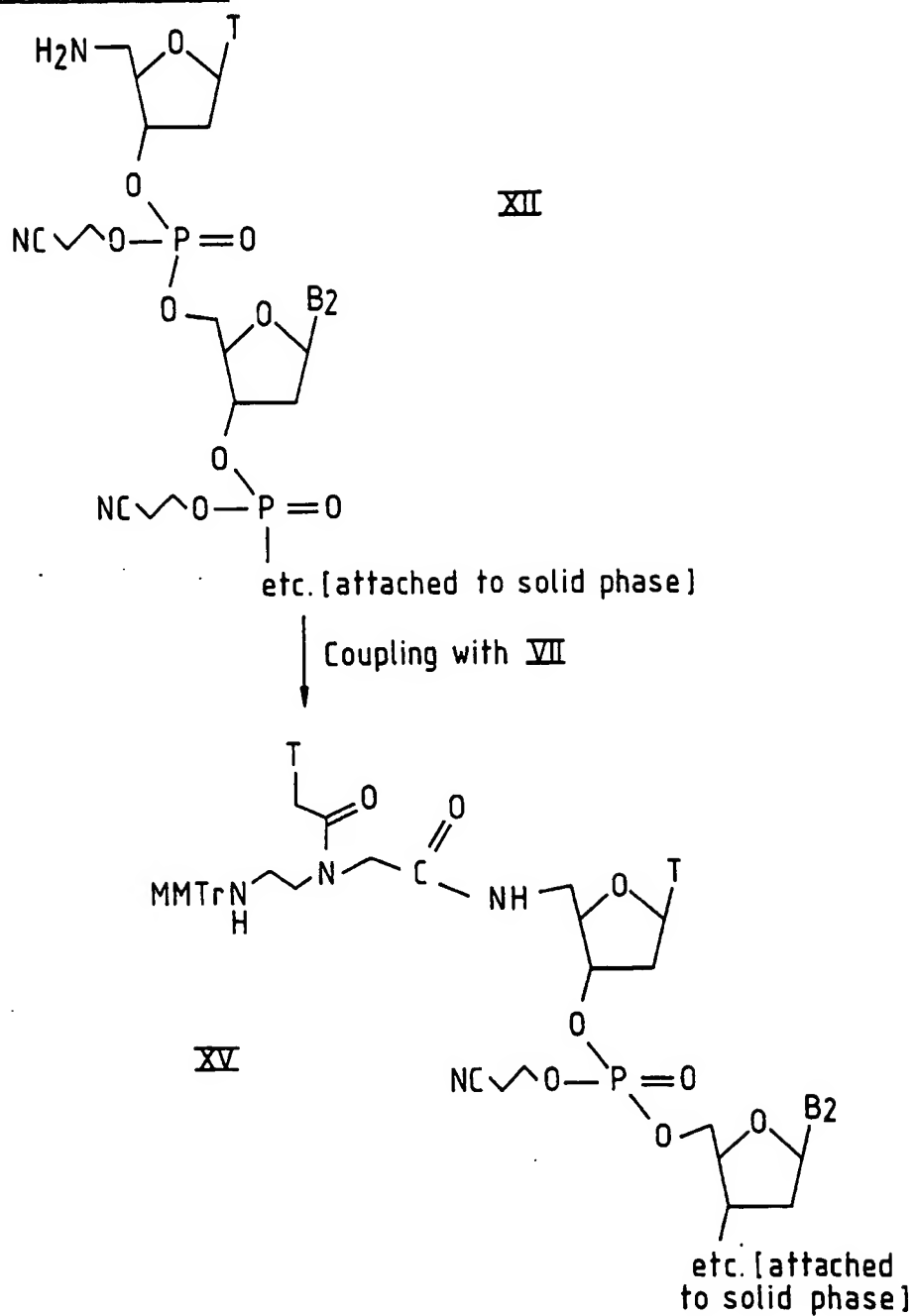
5'-N-(4-Methoxytrityl)amino-5'-deoxythymidine (0.44g, 0.85mmoles) and diisopropylammonium tetrazolide (30mg, 0.25eq.) were coevaporated with pyridine (2x5ml) then MeCN (5ml). The residue was dissolved in MeCN (5ml) then 2-cyanoethyl-N,N,N',N'-tetraisopropylphosphorodiamidite (0.35ml, 1.4eq.) added and the reaction mixture stirred at room temperature for 16h. DCM (50ml) was added and the solution washed with NaHCO₃ (0.5M, 20ml), dried (MgSO₄) then evaporated to an oil which was purified by SiO₂ chromatography (100% EtOAc, 0.30g, 49%). R_f=0.77 (100% EtOAc).

Reaction Scheme 1

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Reaction Scheme 2

- 21 -

Reaction Scheme 3

Example 6Solid Phase Synthesis of PNA/DNA Chimeric Compounds

5 Chimeric molecules were made by initial
synthesis of the DNA moiety using standard
phosphoramidite chemistry and standard automated
synthesis on an ABI 394 synthesiser. The final DNA
monomer addition was a modified nucleoside
10 phosphoramidite; a 5'-amino-5'-2'-dideoxy nucleoside
derivative, protected at the 5'-end with a
monomethoxytrityl group (XI in Reaction Scheme 2). This
was deprotected at the 5'-end with trichloroacetic acid
in the standard way and the solid phase bound
15 oligonucleotide was then ready for the addition of the
PNA monomers. The PNA section of the molecule was then
added in a stepwise fashion using a manual solid phase
methodology.

20 Standard procedure for manual solid phase coupling

Reaction Scheme 3 shows coupling of support bound DNA
with PNA.

The trityl protecting group of the support
25 bound DNA was cleaved using the appropriate cycle on an
ABI 380B DNA synthesiser giving solid phase DNA with a
5'-amino group ready for a PNA coupling reaction XII.
The PNA monomer VII (which is protected with a
monomethoxy trityl group at the amino terminus) 280 mg
30 was dissolved in DMF/Pyridine (1:1 v/v 4.4ml) to yield
a 0.1M solution. An aliquot (0.1ml) was added to the
peptide coupling reagent HBTU (18mg, 5eq.), and DECHA
(20µl, 10 eq.) then manually passed repeatedly through
the column containing the supported DNA for 1h. The
35 coupling procedure was then repeated, to improve yield.
The PNA monomethoxytrityl group was then removed on the

- 23 -

solid phase using trichloroacetic acid and the next PNA coupling reaction was carried out as previously.

When the desired number of monomers had been added, the molecule was cleaved from the solid support using a standard cycle on the DNA synthesiser and the protecting groups removed by heating in NH_3 at 55°C for 5h. The product was purified by reverse phase HPLC. HPLC purification was carried out on a Gilson 303 system using a reverse phase octadecyl stationary phase. The PNA/DNA molecules were purified "trityl off" and the buffer system employed was; $\text{A}=\text{NH}_4\text{OAc}$ (0.1M), $\text{B}=\text{MeCN}$ (30% v/v)/ NH_4OAc (0.1M). The flow rate was 3ml/min, detector 280nm.

The PNA/DNA chimeric molecules in Table 1 below were synthesised according to the above method. PNA moieties appear in lower case and DNA bases in upper case. Equivalent DNA sequences with standard T nucleotides were also synthesised. Figures 1 to 5 show compared HPLC data for the PNA/DNA and DNA sequences. Clear peaks were obtained and in all cases mobility of PNA/DNA molecules was retarded compared to DNA molecules of equivalent length and sequence.

Table 1

SIZE	SEQUENCE	FIGURE NO.
15-mer	tttTAGAGTGTGTT	1b
12-mer	tttTAGAGTGT	2b
9-mer	tttTAGAGT	3b
7-mer	tttTAGA	4b
5-mer	tttTA	5b

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Example 7PNA-DNA Primer Extension Assay5 Annealing Conditions

The 18-mer DNA template comprised the following sequence:

10 5'-CTGAACAACACTCTAAAA-3'

The PNA-DNA primers were synthesised as described in example 6 and shown in Table 1 above.

15 The control primers comprised DNA only, having base sequences identical to the PNA-DNA primers described.

20 Annealing reactions were carried out in a total volume of 10 μ l in the presence of half strength Klenow buffer (1X Klenow buffer contains: 50mM Tris-HCl (pH7.5), 5mM MgCl₂, 5mM β -Mercaptoethanol). The concentrations of primer and template were 5 and 1 pmol respectively.

25 Annealing was performed by boiling the reaction mixtures for 3 minutes and then leaving the samples at room temperature for 30 minutes.

Primer Extension Assay30 The extension reactions contained the contents of the annealing reactions described above (10 μ l each) together with 20 μ M dGTP, 20 μ M dATP, 20 μ M dTTP, 2.5 μ Ci α -³²PdCTP (3000Ci/mmol, Amersham), 1.5X Klenow buffer, Klenow or exonuclease free Klenow (United States
35 Biochemical) at 1 unit per reaction in a total volume of

- 25 -

20µl. (Klenow is a fragment of the enzyme DNA polymerase 1).

5mM d ATP α-S was included in reactions containing

5 Klenow enzyme in order to prevent the 3'-5' exonuclease activity of the said enzyme.

Reactions were incubated at 37°C for up to 30 minutes.

Reaction products were analysed by polycrylamide gel electrophoresis or TLC as described below.

10

Thin Layer Chromatography (TLC) Analysis System

2µl of each reaction mix was added to 3µl of stop

solution (95% formamide, 20mM EDTA, 0.05% bromophenol

15 blue and xylene cyanol FF). 1.0µl of each stopped reaction was then spotted onto a Schleicher and Schuell PEI-cellulose TLC plate 20mm up from the base of the plate. The plates were developed in a tank containing

1M KH₂PO₄. The plates were allowed to run until the

20 solvent front had reached 10mm from the top of the plate. The plates dried and analysed on a Raytest TLC scanner. Labelled reaction products stayed on or close to the origin whereas the [α³²P] dNTP or excised [α³²P] dNMP moved up the plate according to the base used and

25 the number of phosphates.

Denaturing Polyacrylamide gel electrophoresis (PAGE)

Page ITM Sequencing Gel Mix (19:1) from Boehringer

30 Mannheim Corporation was diluted according as described by the manufacturer to give an 18% polyacrylamide gel mix containing 7M urea. 70ml of this mix was taken and polymerised by the addition of 22µl TEMED and 650µl 10% ammonium persulphate. This was used to pour a standard

35 0.4mm sequencing gel. The gel was run in 1x TBE buffer.

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Gels were pre-run for 30 minutes at 1.5kV. Standard sharks tooth combs were used and 2.5 μ l of each sample containing 5000cpm/ μ l in 95% formamide, 20mM EDTA, 0.05% bromophenol blue and xylene cyanol FF was loaded per well. Gels were run for 2 hours at 2.8kV. Gels were fixed in 10% acetic acid: 10% methanol and dried onto Whatman 3MM paper. The dried gel was then exposed to β max film (Amersham).

10

Results

Control experiments were set up using the DNA primers and 18mer template. The DNA oligo primer sizes were 4mer, 5mer, 7mer, 9mer, 12mer, & 15mer respectively. Only the 9, 12, and 15mer gave efficient priming of the template as followed by the incorporation of [α^{32} P]dNTP on TLC plates. The 7mer did prime the template allowing extension but to a lesser degree than the larger DNA oligo's. This was true when both Klenow and exonuclease free Klenow enzymes were used respectively.

Further experiments used the PNA-DNA chimeric primers to prime the template, again priming of the template was followed by the incorporation of [α^{32} P]dNTP when analysed byTLC. Only the PNA-DNA 5mer failed to show any incorporation of the radiolabel. All the other four chimeras showed efficient incorporation of the radiolabel in the presence of either Klenow and dATP α S or exonuclease free Klenow.

The mobility of the labelled reaction products from the PNA-DNA chimera was significantly retarded and ran at a higher molecular weight compared to the corresponding DNA primer reaction products in polyacrylamide gel electrophoresis.

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Competition extension reactions involving the equivalent chimera and DNA primers in the presence of the template resulted in the chimera being preferentially radiolabelled and no radiolabelled DNA product was produced.

Abbreviations

	DCM	dichloromethane
10	DCC	dicyclohexylcarbodiimide
	DCU	dicyclohexylurea
	DECHA	diethylcyclohexylamine
	DMF	dimethylformamide
	EtOAc	ethyl acetate
15	HBTU	hydroxybenzotriazolyltetramethyluroniumhexafluorophosphate
	MMTr	p-Anisoyldiphenylmethane
	HPLC	High Pressure Liquid Chromatography

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CLAIMS

1. A chimeric nucleic acid/nucleic acid analogue
5 molecule comprising a first portion, said first portion comprising a nonstandard backbone oligonucleotide having at least one amide linkage and a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor
10 for the formation of a phosphodiester bond.
2. A chimeric molecule as claimed in claim 1, wherein the second portion comprises at least one normally backbone nucleotide and has a 3' end.
3. A chimeric molecule as claimed in claim 1 or
15 claim 2, wherein the molecule is labelled.
4. A chimeric molecule as claimed in any one of claims 1 to 3, wherein the nonstandard backbone oligonucleotide has a polyamide backbone.
5. A chimeric molecule as claimed in claim 4,
20 wherein the nonstandard backbone oligonucleotide is PNA.
6. A method of performing a primer extension reaction by the use of
 - a) a target nucleic acid
 - 25 b) a primer which is a chimeric nucleic acid/nucleic acid analogue molecule comprising a first portion, said first portion comprising a nonstandard backbone oligonucleotide and a second portion, said second portion comprising an acceptor end which is a
30 chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond, said chimeric molecule being capable of hybridising to part of the target
 - c) a supply of nucleotides
 - 35 which method comprises mixing reagents a), b)

- 29 -

and c) in the presence of a chain extension enzyme under conditions to allow the chimeric molecule to hybridise to the target and extension of the chimeric molecule at the acceptor end to occur, giving an extension product.

7. A method as claimed in claim 6, wherein at least one of reagents a), b) and c) is labelled.

8. A method of amplifying a target nucleic acid by an amplification reaction, wherein at least one primer is used which is a chimeric nucleic acid/nucleic acid analogue molecule comprising a first portion, said first portion comprising a nonstandard backboned oligonucleotide and a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond.

9. A method as claimed in claim 8, wherein the amplification reaction is the polymerase chain reaction.

10. A method of performing a chain termination reaction by the use of

- a) a target nucleic acid
- b) a primer which is a a chimeric nucleic acid/nucleic acid analogue molecule comprising a first portion, said first portion comprising a nonstandard backboned oligonucleotide and a second portion, said second portion comprising an acceptor end which is a chemical functionality capable of acting as acceptor for the formation of a phosphodiester bond, said chimeric molecule being capable of hybridising to part of the target
- c) a supply of nucleotides
- d) a chain termination agent

which method comprises mixing reagents a), b) c) and d) in the presence of a chain extension enzyme under conditions to allow the chimeric molecule to

- 30 -

hybridise to the target and extension of the chimeric molecule at the acceptor end to occur, so as to produce terminated extension products, which terminated extension products are separated to allow
5 part of the nucleotide sequence of the target nucleic acid to be determined.

11. A method as claimed in claim 10, wherein at least one of the reagents b), c) and d) is labelled.

12. A method of determining the nucleotide
10 sequence of a target nucleic acid, which method comprises performing the method of claim 10 or claim 11 using a chain termination agent for each of the four different nucleotides such that the nucleotide sequence of the target may be determined.

15 13. A method as claimed in any one of claims 6 to 12, wherein the target nucleic acid is present in a double-stranded nucleic acid.

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Fig.1a.

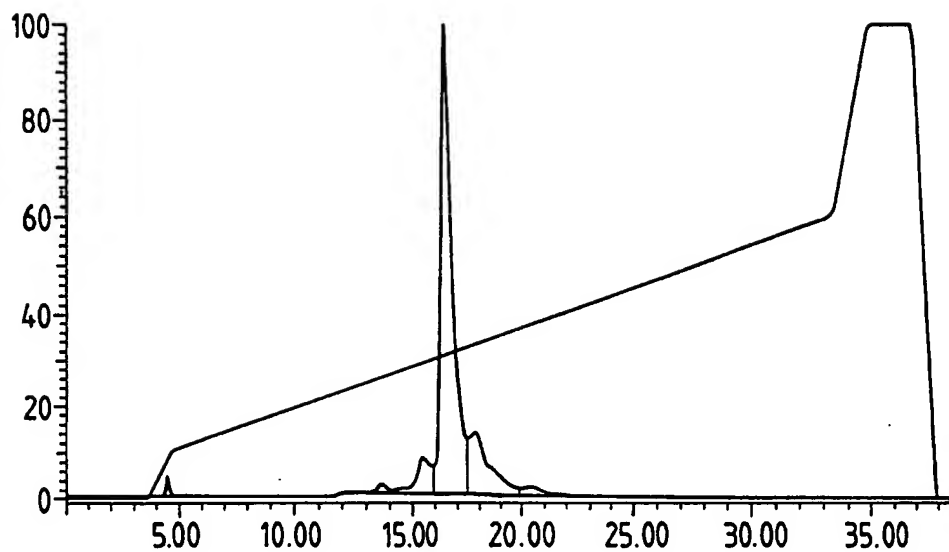
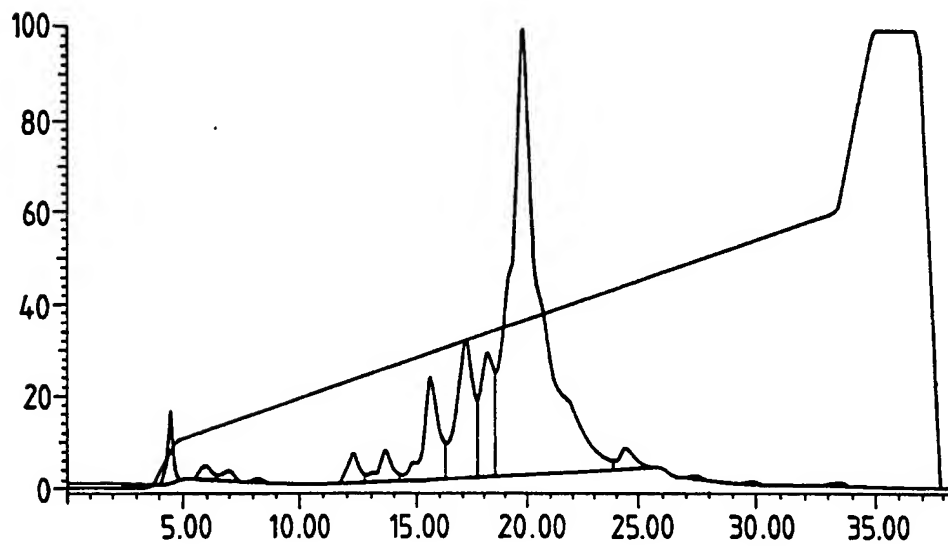


Fig.1b.



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Fig.2a.

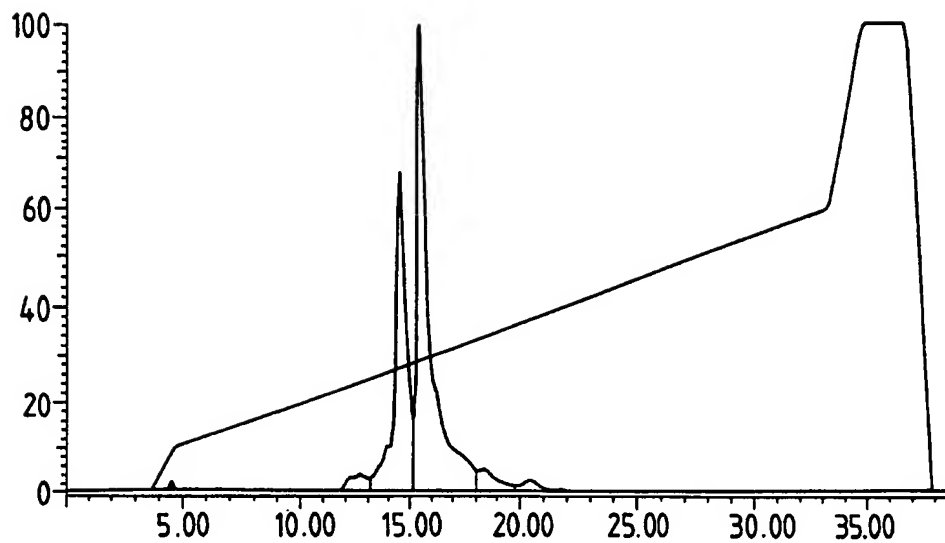
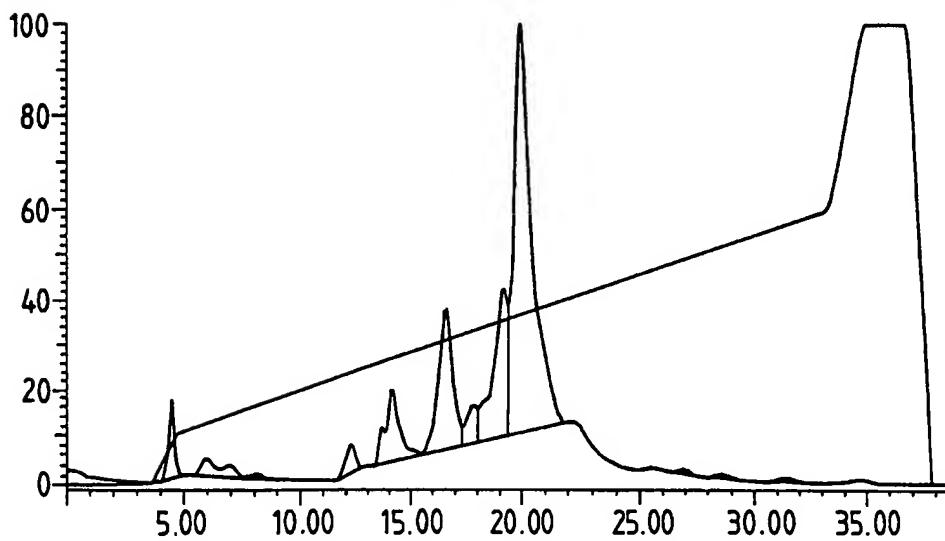


Fig.2b.



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Fig.3a.

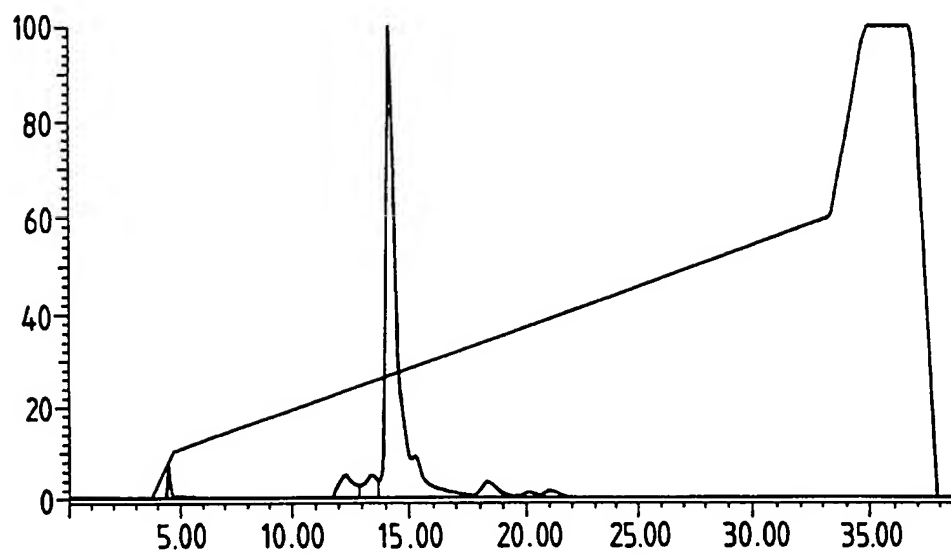
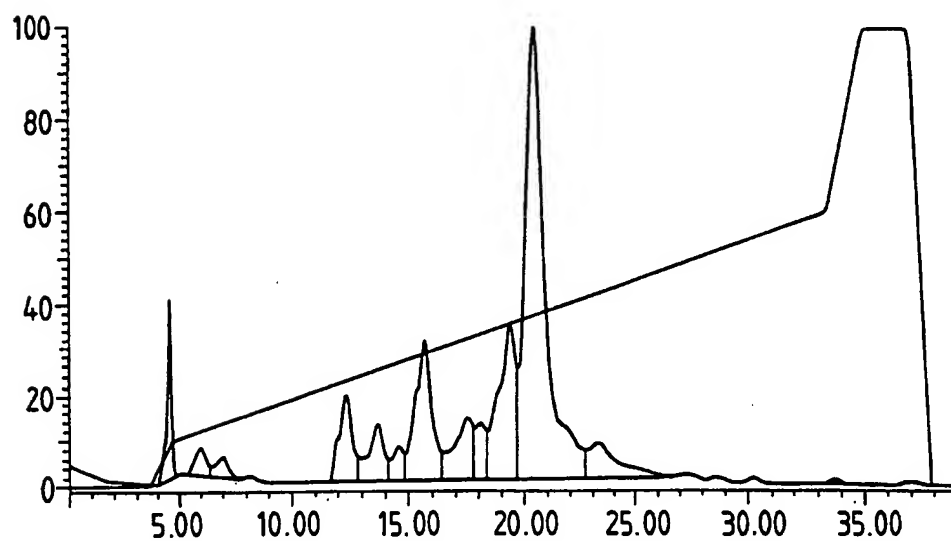


Fig.3b.



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Fig.4a.

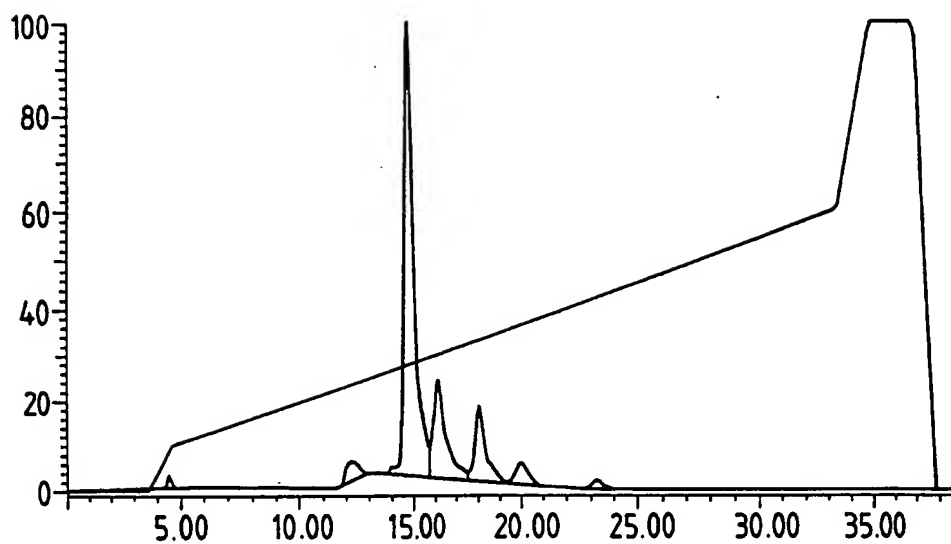
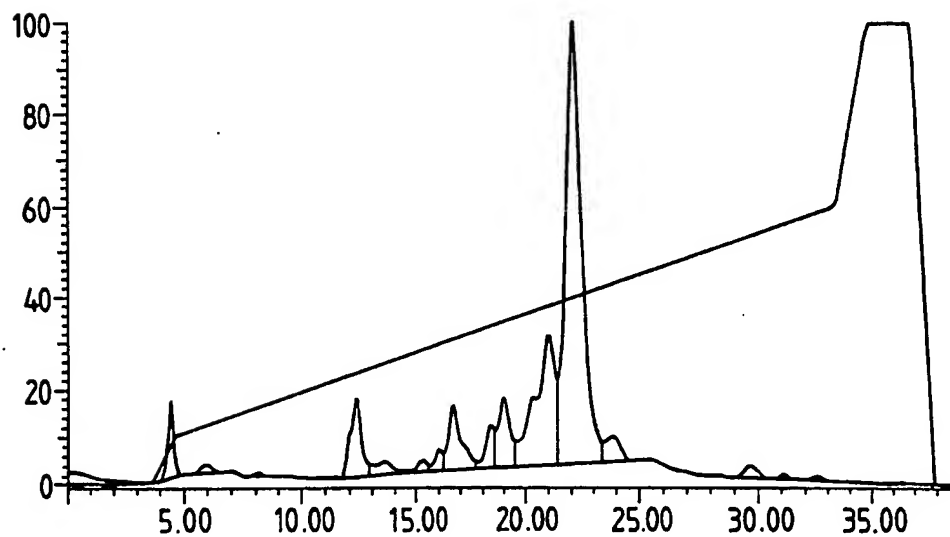


Fig.4b.



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Fig.5a.

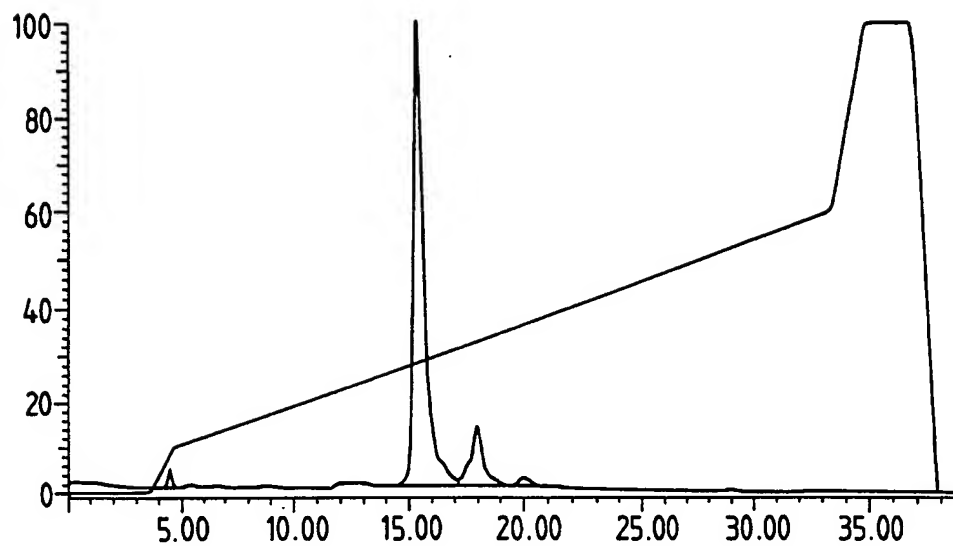
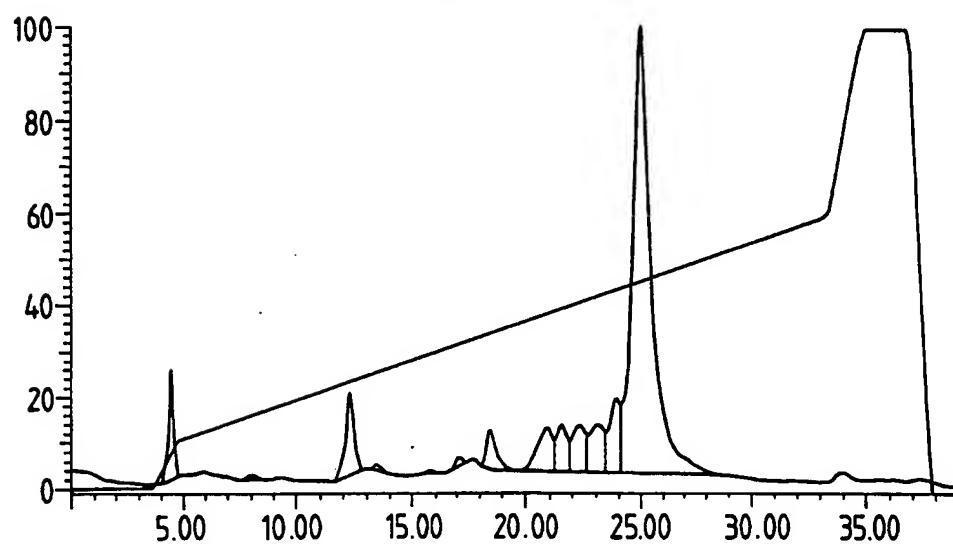


Fig.5b.



INTERNATIONAL SEARCH REPORT

Inter. Appl. No.
PCT/GB 94/02053

A. CLASSIFICATION OF SUBJECT MATTER
IPC 6 C07H21/00 C12Q1/68

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 C07H C12Q

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	EP,A,0 411 186 (ABBOTT LABORATORIES) 6 February 1991 see column 3, line 39 - line 53 see column 6, line 40 - line 46 ---	1-3
X	NUCLEIC ACIDS RESEARCH., vol.20, no.14, 25 July 1992, ARLINGTON, VIRGINIA US pages 3551 - 3554 A.SKERRA see the whole document ---	1-3,6,8
X	EP,A,0 546 761 (BECTON, DICKINSON & COMPANY) 16 June 1993 see page 7, line 9 - line 12 ---	1-3,6-9
-/--		

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

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Date of the actual completion of the international search

25 November 1994

Date of mailing of the international search report

21. 12 94

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INTERNATIONAL SEARCH REPORT

Inter. Application No
PCT/GB 94/02053

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT		
Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	DE,A,41 10 085 (BOEHRINGER INGELHEIM INTERNATIONAL GMBH) 1 October 1992 see page 3, line 20 - line 30 see page 3, line 41 - line 43 ---	1-3,8,9
X	WO,A,92 20703 (O.BUCHARDT ET AL.) 26 November 1992 cited in the application * whole document, especially page 6, lines 23-27 and page 20, lines 35-36 * ---	1-5
A	EP,A,0 224 126 (THE UNIVERSITY OF CALGARY) 3 June 1987 see the whole document ---	10-13
A	TRENDS IN BIOTECHNOLOGY, vol.11, September 1993, CAMBRIDGE GB pages 384 - 386 O.BUCHARDT ET AL. see the whole document -----	1-13

INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/GB 94/02053

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WO-A-9220703	26-11-92	AU-A- 1880692 AU-A- 1884392 CA-A- 2109320 WO-A- 9220702 EP-A- 0586474 EP-A- 0586618 JP-T- 6509063 JP-T- 6506945 NO-A- 934122 NO-A- 934235	30-12-92 30-12-92 25-11-92 26-11-92 16-03-94 16-03-94 13-10-94 04-08-94 11-01-94 20-01-94
EP-A-0224126	03-06-87	JP-A- 63219380	13-09-88